

1631

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE



In re patent application of
KNAPPIK, Achim *et al.*

Application No.: 09/490,064

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For: PROTEIN/ (POLY) PEPTIDE LIBRARIES

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Group Art Unit: 1631

Examiner: M. Zeman

AMENDMENT ACCOMPANYING FORMAL DRAWINGS

Commissioner for Patents
Washington, D.C. 20231

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TECH CENTER 1600/2900

Sir:

Applicants respectfully request that the following amendments be entered into
the application:

IN THE SPECIFICATION:

Page 18, line 4, paragraph 2: Please amend as follows:

E1
Figs. 2A-2G: Alignment of consensus sequences designed for each subgroup
(amino acid residues are shown with their standard one-letter abbreviation). (2A-2B)
kappa sequences, (2C-2D) lambda sequences and (2E-2G), heavy chain sequences.
The positions are numbered according to Kabat (1991). In order to maximize homology
in the alignment, gaps (-) have been introduced in the sequence at certain positions.

E2
[Page 18, line 10, paragraph 3: Please amend as follows:]

Figs. 3A-3K: Gene sequences of the synthetic V kappa consensus genes. The
corresponding amino acid sequences (see Figs. 2A-2B) as well as the unique cleavage
sites are also shown.

E3
[Page 18, line 13, paragraph 4: Please amend as follows:]

Figs. 4A-4I: Gene sequences of the synthetic V lambda consensus genes.
The corresponding amino acid sequences (see Figs. 2C-2D) as well as the unique
cleavage sites are also shown.

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